SEQUENCE LISTING

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<120> Improved Method of Purifying TFPI and TFPI Analogs

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<150> US 60/494,546
<151> 2003-08-13

<150> US 60/509,277
<151> 2003-10-08

<150> US 60/512,199
<151> 2003-10-20

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Gly Pro Cys Lys Ala Ile Met Lys Arg Phe Phe Phe Asn Ile Phe Thr
        35                  40                  45
Arg Gln Cys Glu Glu Phe Ile Tyr Gly Gly Cys Glu Gly Asn Gln Asn
    50                  55                  60
Arg Phe Glu Ser Leu Glu Glu Cys Lys Lys Met Cys Thr Arg Asp Asn
65                  70                  75                  80
Ala Asn Arg Ile Ile Lys Thr Thr Leu Gln Gln Glu Lys Pro Asp Phe
                85                  90                  95
Cys Phe Leu Glu Glu Asp Pro Gly Ile Cys Arg Gly Tyr Ile Thr Arg
            100                 105                 110
Tyr Phe Tyr Asn Asn Gln Thr Lys Gln Cys Glu Arg Phe Lys Tyr Gly
        115                 120                 125
Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu Glu Glu Cys Lys
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Asn Ile Cys Glu Asp Gly Pro Asn Gly Phe Gln Val Asp Asn Tyr Gly
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Thr Gln Leu Asn Ala Val Asn Asn Ser Leu Thr Pro Gln Ser Thr Lys
                165                 170                 175
Val Pro Ser Leu Phe Glu Phe His Gly Pro Ser Trp Cys Leu Thr Pro
            180                 185                 190
Ala Asp Arg Gly Leu Cys Arg Ala Asn Glu Asn Arg Phe Tyr Tyr Asn
        195                 200                 205
Ser Val Ile Gly Lys Cys Arg Pro Phe Lys Tyr Ser Gly Cys Gly Gly
    210                 215                 220
Asn Glu Asn Asn Phe Thr Ser Lys Gln Glu Cys Leu Arg Ala Cys Lys
225                 230                 235                 240
Lys Gly Phe Ile Gln Arg Ile Ser Lys Gly Gly Leu Ile Lys Thr Lys
                245                 250                 255
Arg Lys Arg Lys Lys Gln Arg Val Lys Ile Ala Tyr Glu Glu Ile Phe
            260                 265                 270
Val Lys Asn Met
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Asp Glu Glu His Thr Ile Ile Thr
 1               5

<210> 3
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 1               5

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            20                  25                  30
Cys Arg Pro Phe Lys Tyr Ser Gly Cys Gly Gly Asn Glu Asn Asn Phe
        35                  40                  45
Thr Ser Lys Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly Phe Ile Gln
    50                  55                  60
Arg Ile Ser Lys Gly Gly Leu Ile Lys Thr Lys Arg Lys Arg Lys Lys
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20 25 30
Cys Arg Pro Phe Lys Tyr Ser Gly Cys Gly Gly Asn Glu Asn Asn Phe
35 40 45
Thr Ser Lys Gln Glu Cys Leu Arg Ala Cys Lys Lys Gly Phe Ile Gln
50 55 60
Arg Ile Ser Lys Gly Gly Leu Ile Lys Thr Lys Arg Lys Arg Lys Lys
65 70 75 80
Gln Arg Val Lys Ile Ala Tyr
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Gln Ser Thr Lys Val Pro Ser Leu Phe
20 25

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35 40 45
Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu Glu Glu
50 55 60
Cys Lys Asn Ile Cys Glu Asp Gly Pro Asn Gly Phe Gln Val
65 70 75

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Lys Ala Asp Asp Gly Pro Cys Lys Ala Ile Met Lys Arg Phe Phe Phe
            20   25   30
Asn Ile Phe Thr Arg Gln Cys Glu Glu Phe Ile Tyr Gly Gly Cys Glu
        35   40   45
Gly Asn Gln Asn Arg Phe Glu Ser Leu Glu Glu Cys Lys Lys Met Cys
    50   55   60
Thr Arg
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Lys Gln Cys Glu Arg Phe
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Lys Ala Ile Met Lys Arg
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Lys Gln Glu Cys Leu Arg Ala
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 1   5

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Lys Cys Arg Pro Phe Lys Tyr
 1   5

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Lys Lys Gly Phe Ile Gln Arg Ile
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Lys Phe Glu Ser Leu Glu Glu Cys Lys Lys Met
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Arg Phe Glu Ser Leu Glu Glu Cys Lys Lys
1 5 10

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<400> 24
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<400> 25
Lys Leu Met His Ser Phe Cys Ala Phe Lys Ala
 1   5   10

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<400> 26
Lys Ile Ala Tyr Glu Glu Ile Phe Val Lys Asn
 1   5   10

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<400> 27
Lys Asn Ile Cys Glu Asp Gly Pro Asn Gly Phe Gln Val Asp Asn Tyr
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Lys Val

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Lys Tyr Gly Gly Cys Leu Gly Asn Met Asn Asn Phe Glu Thr Leu Glu
 1   5   10   15
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Ala Asp Ser Glu Glu Asp Glu Glu His Thr Ile Ile Thr Asp Thr Glu
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Leu Pro Pro Leu Lys Leu
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Asp Pro Gly Ile Cys Arg Gly
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<400> 31
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Glu Cys Lys Asn Ile Cys Glu Asp Gly Pro Asn Gly Phe Gln Val Asp
            20   25   30
Asn Tyr Gly Thr Gln Leu Asn Ala Val Asn Asn Ser Leu Thr Pro Gln
        35   40   45
Ser Thr Lys Val
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Lys Val Pro Ser Leu Phe Glu Phe His Gly Pro Ser Trp Cys Leu Thr
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<400> 33
Arg Phe Phe Phe Asn Ile Phe Thr Arg Gln
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Thr Thr Leu Gln Gln Glu Lys Pro Asp Phe Cys Phe Nva Glu Glu Asp
 1   5   10   15
Pro Gly Ile Cys Arg
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 1   5   10   15
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 1   5   10   15
Ala Asp Arg

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Ala Asp Arg

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catgaagcgc ttcttcttca acatcttcac tcgtcagtgc gaagaattta tatatggggg 180
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